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Corrigendum

Corrigendum to "Rpa43 and its partners in the yeast RNA polymerase I transcription complex" [FEBS Lett. 585 (21) (2011) 3355–3359]

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Two tables (Tables 1 and 2) are missing in the original text.

Part 3.2:

Conversely, N-terminal deletions (rpa43-35,326, rpa43-35,281) had a wild-type growth, except for their dependency on Rpa14. They were not synthetic-lethal with rpa12D, rpa34D or rpa49D but suppressed the cold-sensitive defect of rpa49D (Table 2).

Table 1
Conservation of Pol I subunits and initiation factors. Data based on a Psi-blast survey of eukaryotic genomes (http://blast.ncbi.nlm.nih.gov/Blast.cgi). Rpa43 and Rpa14 alignments are provided as Supplementary data S2.

Lineages	Organisms	Rpa43	Rpa14	Rpa49	Rpa34	Rrn3 TIF-1A	Rrn7 TIF-1B	Rrn6 TIF-1C
Unikonts								
Fungi	*S. cerevisiae*	+	+	+	+	+	+	+
Animals	*H. sapiens*	+		+	+	+	+	+
Amoebozoa	*D. discoideum*	+		+	+	+	+	+
Plantae								
Plants	*A. thaliana*	+		+		+	+	+
Green algae	*C. reinhardtii*	+		+		+	+	+
Others								
Oomycetes	*P. infestans*	+		+		+	+	+
Apicomplexans	*T. gondii*	+				+		
Ciliates	*T. thermophila*	+		+		+		
Brown algae	*E. siliculosus*	+		+		+		
Diplomonads	*G. lamblia*	+		+		+		
Trichomonas	*T. vaginalis*	+				+		

Table 2
Synthetic growth defect between *rpa43* alleles and deletions of non-essential Pol I subunits (Rpa12, Rpa14, Rpa34, Rpa49) or of the Pol I-associated factors Hmo1 and Top1. A suppression of the cs defect of *rpa49Δ* is underscored in blue.

RPA43	WT	*rpa14Δ*	*rpa12Δ*	*rpa34Δ*	*rpa49Δ*	*top1Δ*	*hmo1Δ*
WT	++	++	+,ts	++	cs,+, ts	+	cs,+
chim1	++	-	-	ts	-	+	cs,+
chim2	+, ts	-	-	-	-	+, ts	cs,(+),ts
1, 235	++	-	-	-	-	+, (ts)	cs,+
1, 265	++	++	+,ts	++	cs,+, ts	+	cs,+
1, 281	++	++	+,ts	++	cs,+, ts	+	cs,+
35, 326	++	-	+,ts	++	+,ts	+	cs,+
35, 281	++	ts	+,ts	++	+,ts *	+	cs,+

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